WA

PCT10

RAW SEQUENCE LISTING DATE: 02/21/2003
PATENT APPLICATION: **US/10/089,503A** TIME: 10:00:55

Input Set : **N:\Crf4\01302003\J089503.raw**
Output Set: **N:\CRF4\02212003\J089503A.raw**

<110> APPLICANT: Universitatsklinikum Freiburg
<120> TITLE OF INVENTION: THE PRV-1 GENE AND USE THEREOF
<130> FILE REFERENCE: E980930
<140> CURRENT APPLICATION NUMBER: US/10/089,503A
<141> CURRENT FILING DATE: 2002-09-20
<150> PRIOR APPLICATION NUMBER: DE 199 47 010.3
<151> PRIOR FILING DATE: 1999-09-30
<160> NUMBER OF SEQ ID NOS: 10
<170> SOFTWARE: PADAT Sequenzmodul, Version 1.0

ENTERED

<210> SEQ ID NO: 1
<211> LENGTH: 1600
<212> TYPE: DNA
<213> ORGANISM: homo sapiens
<220> FEATURE:
<223> OTHER INFORMATION:

W--> **<400> 1**

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ttcagcatgt gtggaaggtg tccgacctgc cccggcaatg gacccctaag aacaccagct 180
gcgacagcgg cttggggtgc caggacacgt tgatgctcat tgagagcgga ccccaagtga 240
gcctggtgct ctccaagggc tgcacggagg ccaaggacca ggagccccgc gtcactgagc 300
accggatggg ccccggcctc tccctgatct cctacacctt cgtgtgccgc caggaggact 360
tctgcaacaa cctcgttaac tccctcccgc tttgggcccc acagccccca gcagacccag 420
gatccttgag gtgcccagtc tgcttgtcta tggaaggctg tctggagggg acaacagaag 480
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atgggacaca ggaaattggg cccgtgggta tgactgagaa ctgcaatagg aaagattttc 660
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attggaccac atcgaatacc gagatgtgcg aggtggggca ggtgtgtcag gagacgctgc 780
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Input Set : **N:\Crf4\01302003\J089503.raw**
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<210> SEQ ID NO: 2
<211> LENGTH: 437
<212> TYPE: PRT
<213> ORGANISM: homo sapiens
<400> SEQUENCE: 2

```
Met Ser Ala Val Leu Leu Leu Ala Leu Leu Gly Phe Ile Leu Pro Leu
 1               5                  10                  15
Pro Gly Val Gln Ala Leu Leu Cys Gln Phe Gly Thr Val Gln His Val
            20                  25                  30
Trp Lys Val Ser Asp Leu Pro Arg Gln Trp Thr Pro Lys Asn Thr Ser
        35                  40                  45
Cys Asp Ser Gly Leu Gly Cys Gln Asp Thr Leu Met Leu Ile Glu Ser
    50                  55                  60
Gly Pro Gln Val Ser Leu Val Leu Ser Lys Gly Cys Thr Glu Ala Lys
65                  70                  75                  80
Asp Gln Glu Pro Arg Val Thr Glu His Arg Met Gly Pro Gly Leu Ser
                85                  90                  95
Leu Ile Ser Tyr Thr Phe Val Cys Arg Gln Glu Asp Phe Cys Asn Asn
            100                 105                 110
Leu Val Asn Ser Leu Pro Leu Trp Ala Pro Gln Pro Pro Ala Asp Pro
        115                 120                 125
Gly Ser Leu Arg Cys Pro Val Cys Leu Ser Met Glu Gly Cys Leu Glu
    130                 135                 140
Gly Thr Thr Glu Glu Ile Cys Pro Lys Gly Thr Thr His Cys Tyr Asp
145                 150                 155                 160
Gly Leu Leu Arg Leu Arg Gly Gly Gly Ile Phe Ser Asn Leu Arg Val
                165                 170                 175
Gln Gly Cys Met Pro Gln Pro Gly Cys Asn Leu Leu Asn Gly Thr Gln
            180                 185                 190
Glu Ile Gly Pro Val Gly Met Thr Glu Asn Cys Asn Arg Lys Asp Phe
        195                 200                 205
Leu Thr Cys His Arg Gly Thr Thr Ile Met Thr His Gly Asn Leu Ala
    210                 215                 220
Gln Glu Pro Thr Asp Trp Thr Thr Ser Asn Thr Glu Met Cys Glu Val
225                 230                 235                 240
Gly Gln Val Cys Gln Glu Thr Leu Leu Leu Ile Asp Val Gly Leu Thr
                245                 250                 255
Ser Thr Leu Val Gly Thr Lys Gly Cys Ser Thr Val Gly Ala Gln Asn
            260                 265                 270
Ser Gln Lys Thr Thr Ile His Ser Ala Pro Pro Gly Val Leu Val Ala
        275                 280                 285
Ser Tyr Thr His Phe Cys Ser Ser Asp Leu Cys Asn Ser Ala Ser Ser
    290                 295                 300
Ser Ser Val Leu Leu Asn Ser Leu Pro Pro Gln Ala Ala Pro Val Pro
305                 310                 315                 320
Gly Asp Arg Gln Cys Pro Thr Cys Val Gln Pro Leu Gly Thr Cys Ser
                325                 330                 335
Ser Gly Ser Pro Arg Met Thr Cys Pro Arg Gly Ala Thr His Cys Tyr
            340                 345                 350
```

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```
Asp Gly Tyr Ile His Leu Ser Gly Gly Gly Leu Ser Thr Lys Met Ser
        355                 360                 365
Ile Gln Gly Cys Val Ala Gln Pro Ser Ser Phe Leu Leu Asn His Thr
    370                 375                 380
Arg Gln Ile Gly Ile Phe Ser Ala Arg Glu Lys Arg Asp Val Gln Pro
385                 390                 395                 400
Pro Ala Ser Gln His Glu Gly Gly Gly Ala Glu Gly Leu Glu Ser Leu
                405                 410                 415
Thr Trp Gly Val Gly Leu Ala Leu Ala Pro Ala Leu Trp Trp Gly Val
            420                 425                 430
Val Cys Pro Ser Cys
        435
```

<210> SEQ ID NO: 3
<211> LENGTH: 24
<212> TYPE: RNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: 5'-end of PRV-1-sequence
<400> SEQUENCE: 3
aaaagcagaa agagauuacc agcc 24

<210> SEQ ID NO: 4
<211> LENGTH: 24
<212> TYPE: RNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Antisense-Molecule
<400> SEQUENCE: 4
ggcugguaau cucuuucugc uuuu 24

<210> SEQ ID NO: 5
<211> LENGTH: 13
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: amino acids 34-46 of PRV-1
<400> SEQUENCE: 5
Lys Val Ser Asp Leu Pro Arg Gln Trp Thr Pro Lys Asn
1 5 10

<210> SEQ ID NO: 6
<211> LENGTH: 15
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: amino acids 391-405 of PRV-1
<400> SEQUENCE: 6
Ser Ala Arg Glu Lys Arg Asp Val Gln Pro Pro Ala Ser Gln His
1 5 10 15

<210> SEQ ID NO: 7
<211> LENGTH: 27
<212> TYPE: DNA

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Output Set: **N:\CRF4\02212003\J089503A.raw**

149 <213> ORGANISM: Artificial Sequence
150 <220> FEATURE:
151 <223> OTHER INFORMATION: RT-Primer
152 <400> SEQUENCE: 7
153 attaggttat gaggtcagag ggaggtt 27
155 <210> SEQ ID NO: 8
156 <211> LENGTH: 28
157 <212> TYPE: DNA
158 <213> ORGANISM: Artificial Sequence
159 <220> FEATURE:
160 <223> OTHER INFORMATION: sense-Primer
161 <400> SEQUENCE: 8
162 gcagaaagag attaccagcc acagacgg 28
164 <210> SEQ ID NO: 9
165 <211> LENGTH: 28
166 <212> TYPE: DNA
167 <213> ORGANISM: Artificial Sequence
168 <220> FEATURE:
169 <223> OTHER INFORMATION: antisense-Primer
170 <400> SEQUENCE: 9
171 gaatcgtggg ggtaatagag ttagcagg 28
173 <210> SEQ ID NO: 10
174 <211> LENGTH: 29
175 <212> TYPE: DNA
176 <213> ORGANISM: Artificial Sequence
177 <220> FEATURE:
178 <223> OTHER INFORMATION: probe
179 <400> SEQUENCE: 10
180 ttcttgttga accacaccag acaaatcgg 29

RAW SEQUENCE LISTING ERROR SUMMARY DATE: 02/21/2003
PATENT APPLICATION: **US/10/089,503A** TIME: 10:00:56

Input Set : **N:\Crf4\01302003\J089503.raw**
Output Set: **N:\CRF4\02212003\J089503A.raw**

Invalid Line Length:
The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:1; Line(s) 18,19,20,21,22,23,24,25,26,27,28,29,30,31,32,33,34,35,36,37
Seq#:1; Line(s) 38,39,40,41,42,43,44

VERIFICATION SUMMARY DATE: 02/21/2003
PATENT APPLICATION: **US/10/089,503A** TIME: 10:00:56

Input Set : **N:\Crf4\01302003\J089503.raw**
Output Set: **N:\CRF4\02212003\J089503A.raw**

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